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Errors Corrected by the STIC Systems Branch

1644

Serial Number: 09/483,672A

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H6

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☒ Other: Seq 378 - converted a.a. res  
Seq 525 - aligned a.a. res

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

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## RAW SEQUENCE LISTING

DATE: 07/30/2000

PATENT APPLICATION: US/09/483,672A

TIME: 17:04:25

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07282000\I483672A.raw

3 <110> APPLICANT: Xu, Jiangchun  
4 Dillon, Davin C.  
5 Mitcham, Jennifer L.  
6 Harlocker, Susan Louise  
7 Jiang Yuqui  
8 Reed, Steven G.  
9 Kalos, Michael D.  
10 Fanger, Gary R.  
11 Retter, Marc W.  
12 Solk, John A.  
13 Day, Craig H.  
14 Skeiky, Yasir A.W.  
15 Wang, Aijun  
16 Meagher, Madeleine  
18 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND  
19 DIAGNOSIS OF PROSTATE CANCER  
21 <130> FILE REFERENCE: 210121.42711C11  
C--> 23 <140> CURRENT APPLICATION NUMBER: US/09/483,672A  
24 <141> CURRENT FILING DATE: 2000-01-14  
26 <160> NUMBER OF SEQ ID NOS: 590  
28 <170> SOFTWARE: FastSEQ for Windows Version 3.0  
30 <210> SEQ ID NO: 1  
31 <211> LENGTH: 814  
32 <212> TYPE: DNA  
33 <213> ORGANISM: Homo sapien  
35 <220> FEATURE:  
36 <221> NAME/KEY: misc\_feature  
37 <222> LOCATION: (1)...(814)  
38 <223> OTHER INFORMATION: n = A,T,C or G  
40 <400> SEQUENCE: 1  
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43 ccagggggtc cagtccctct ccttacttca tccccatccc atgccaaagg aagaccctcc 180  
44 ctcccttggt cacagccttc tctaggcttc ccagtgcctc caggacagag tgggttatgt 240  
45 ttccagctcc atccttgctg tgagtgtctg gtgcgttggt cctccagctt ctgctcagtg 300  
46 cttcatggac agtgtccagc acatgtcact ctccactctc tcagtgtgga tccactagtt 360  
47 ctagagcggc cgccaccgcg gtggagctcc agcttttggt ccctttagtg agggttaatt 420  
48 ggcgcgttgg cgtaaatcatg gtcataactg ttctctgtgt gaaattgta tccgctcaca 480  
49 attccacaca acatacagag cggaagcata aagtgtaaag cctgggggtc ctaatgagtg 540  
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W--> 53 actcctcaaa ggnggtatta cggttatccn naaatcnggg gatacccngg aaaaaanttt 780  
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56 <210> SEQ ID NO: 2  
57 <211> LENGTH: 816  
58 <212> TYPE: DNA

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PATENT APPLICATION: US/09/483,672A

TIME: 17:04:25

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07282000\I483672A.raw

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68 ttcatggctg ttggagcaat agaaccocag ttctacgagc tgctgatcaa aggacttgga      120
69 cttaaagtctg atgaacttcc caatcagatg agcatggatg attggccaga aatgaagaag      180
70 aagtttgcag atgtatttgc aaagaagacg aaggcagagt ggtgtcaaat ctttgacggc      240
71 acagatgcct gtgtgactcc ggttctgact tttagaggag ttgttcatca tgatcacaaac      300
72 aaggaaacggg gctcgtttat caccagttag gagcaggacg tgagcccccg cctgcacct      360
73 ctgctgttaa acaccccagc catcccttct ttcaaaaggg atccactagt tctagaagcg      420
74 gccgccacgg cggtggagct ccagcttttg ttccctttag tgagggttaa ttgcgcgctt      480
75 gccgtaatca tggtcatagc tgtttcctgt gtgaaattgt tatccgctca caattccccc      540
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W--> 77 cattaattgc gttgcgctca ctgcccgctt tccagtcggg aaaactgtcg tgcactgcn      660
W--> 78 ttantgaatc ngccaccccc cgggaaaagg cggttgcntt ttgggacctc tccgctttcc      720
W--> 79 tcgctcattg atcctngcnc ccggtcttcg gctgcggnga acggttctact cctcaaaggc      780
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88 <221> NAME/KEY: misc_feature
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90 <223> OTHER INFORMATION: n = A,T,C or G
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94 tctgtctcct cactgggtgat aaacgagccc cgttccttgt tggatcatg atgaacaacc      120
95 tctcaaaaag tcagaacccg agtcacacag gcatctgtgc cgtcaaaagt ttgacaccac      180
96 tctgccttcg tcttctttgc aaatacatct gcaaaattct tcttcatttc tggccaatca      240
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98 tcgtagaact ggggttctat tgctccaaca gccatgaatt ccccatctgc tgcctgttaa      360
99 gtcgtataga aagggtgctc accatccaac atgttctgtc ctgcaggggg ggcccggtag      420
W--> 100 ccaattcgcc ctatantgag tcgtattacg cgcgctcact ggccgtcggt ttacaacgtc      480
101 gtgactggga aaaccctggg cgttaccaac ttaatcgctt tgcagacat ccccttttcg      540
W--> 102 ccagctgggc gtaatanaga aaaggcccgcc accgatcgcc cttccaacag ttgcgcacct      600
W--> 103 gaatgggnaa atgggacccc cctgttaccg cgcattnaac ccccgcnngg tttngttggt      660
W--> 104 acccccacnt nnaccgctta cactttgcca gcgccttanc gcccgctccc tttncctttt      720
W--> 105 cttcccttcc tttcnncncc ctttcccccg gggtttcccc cntcaaaccc cna      773
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108 <211> LENGTH: 828
109 <212> TYPE: DNA
110 <213> ORGANISM: Homo sapien
112 <220> FEATURE:
113 <221> NAME/KEY: misc_feature
114 <222> LOCATION: (1)...(828)
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## RAW SEQUENCE LISTING

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PATENT APPLICATION: US/09/483,672A

TIME: 17:04:25

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07282000\I483672A.raw

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119 aatgggcaga cacagggtga tgccaatggt tctgaaatgg gtataatttc gtcctctcct 120
120 tcggaacact ggtgtgtctt gaagacttct cgtcagttt cagtggaggac acacacaaag 180
121 acgtgggtga ccatgttggt tgtgggtgac agagatggga ggggtggggc ccaccctgga 240
122 agagtggaga gtgacacaag gtggacactc tctacagatc actgaggata agctggagcc 300
W--> 123 acaatgcatg aggcacacac acagcaagga tgacnctgta aacatagccc acgtgtctct 360
W--> 124 gngggcactg ggaagcctan atnaggcctg gagcanaaag aaggggagga tccactagtt 420
W--> 125 ctanagcggc cgccaccgag gtgganctcc ancttttggt cccttttagtg agggttaatt 480
W--> 126 gcgcgcttgg cntaatcatg gtcatanctn tttcctgtgt gaaattgtta tccgctcaca 540
W--> 127 attccacaca acatacganc cggaaacata aantgtaaac ctgggggtgcc taatgantga 600
W--> 128 ctaactcaca ttaattgcgt tgcgctcact gcccgcttcc caatcnggaa acctgtcttg 660
W--> 129 ccncttgcat tnatgaaten gccaaccccc ggggaaaagg gtttgcgttt tgggcgctct 720
W--> 130 tccgcttctc cnetcantta ntccctnnc tccgtcattc cggctgcngc aaaccgggtc 780
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136 <213> ORGANISM: Homo sapien
138 <220> FEATURE:
139 <221> NAME/KEY: misc_feature
140 <222> LOCATION: (1)...(834)
141 <223> OTHER INFORMATION: n = A,T,C or G
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145 agtttttaatt gcatccaaag tactaacaac aactctagca atcaagaatg gcagcatggt 120
146 attttataac aatcaacacc tgtggctttt aaaatttgggt ttccataaga taattttatac 180
147 tgaagtaaat ctagccatgc ttttaaaaaa tgcttttaggt cactccaagc ttggcagtta 240
148 acatttggca taaacaataa taaaacaatc acaatttaac aaataacaaa tacaacattg 300
149 tagggcataa tcataacag tataaggaaa aggtggtagt gttgagtaag cagttattag 360
150 aatagaatac ctggcctct atgcaaatat gtctagacac ttgatttoac tcagccctga 420
151 cattcagttt tcaaatgtag agacaggttc tacagtatca tttacagtt tccaacacat 480
152 tgaaaaaag tagaaaatga tgagttgatt tttattaatg cattacatcc tcaagagtta 540
153 tcaccaaccc ctcagttata aaaaattttc aagttatatt agtcatataa cttggtgtgc 600
154 ttattttaaa ttagtgtctaa atggattaag tgaagacaac aatggtcccc taatgtgatt 660
W--> 155 gatattgggc atttttacca gtttctaaat ctnaacttcc aggcctttga actggaacat 720
W--> 156 tgnatnacag tgttccanag ttncaccta ctggaacatt acagtgtgct tgattcaaaa 780
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160 <211> LENGTH: 818
161 <212> TYPE: DNA
162 <213> ORGANISM: Homo sapien
164 <220> FEATURE:
165 <221> NAME/KEY: misc_feature
166 <222> LOCATION: (1)...(818)
167 <223> OTHER INFORMATION: n = A,T,C or G
169 <400> SEQUENCE: 6
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## RAW SEQUENCE LISTING

DATE: 07/30/2000

PATENT APPLICATION: US/09/483,672A

TIME: 17:04:25

Input Set : A:\Pto.amc

Output Set : N:\CRF3\07282000\I483672A.raw

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172 tgtaaaagtga aatattagtt ggcggatgaa gcagatagtg aggaaagtgg agccaataat 180
173 gacgtgaagt ccttggaagc ctgtggctac aaaaaatggt gagccgtaga tgccgtcggg 240
174 aatgggtgaag ggagactcga agtactctga ggcctttagg agggtaaaat agagaccag 300
175 taaaattgta ataagcagtg cttgaattat ttggtttcgg ttgttttcta ttagactatg 360
176 gtgagctcag gtgattgata ctctgatgc gagtaatacg gatgtgttta ggagtgggac 420
177 ttctagggga tttagcgggg tgatgcctgt tgggggccag tgccctccta gttggggggt 480
178 aggggctagg ctggagtggg aaaaggctca gaaaaatcct gcgaagaaaa aaacttctga 540
179 ggtaataaat aggattatcc cgtatcgaag gcctttttgg acagggtggg tggtgtggcc 600
180 ttggtatgtg ctttctcgtg ttacatcgcg ccatcattgg tataatggtta gtgtgttggg 660
W--> 181 ttantanggc ctantatgaa gaacttttgg antggaatta aatcaatngc ttggccggaa 720
W--> 182 gtcattanga nggctnaaaa ggcctgttta ngggtctggg ctnggtttta cccnaccat 780
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187 <212> TYPE: DNA
188 <213> ORGANISM: Homo sapien
190 <220> FEATURE:
191 <221> NAME/KEY: misc_feature
192 <222> LOCATION: (1)...(817)
193 <223> OTHER INFORMATION: n = A,T,C or G
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198 ggtttgctcc acagatttca gacattgac cgtagtatac ccccggtcgt gtagcgggtga 180
199 aagtgttttg gtttagacgt ccgggaattg catctgtttt taagcctaata gtggggacag 240
W--> 200 ctcatgagtg caagacgtct tgtgatgtaa ttattatacn aatggggggt tcaatcggga 300
201 gtactactcg attgtcaacg tcaaggagtc gcaggtcgcc tggttctagg aataatgggg 360
202 gaagtatgta ggaattgaag attaatccgc cgtagtcggt gttctcctag gttcaatacc 420
203 attggtggcc aattgatttg atggttaagg gagggatcgt tgaactcgtc tgttatgtaa 480
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W--> 205 tcaaacngtc tctanttcct gaaacgtctg aaatgttaat aanaattaan tttngttatt 600
W--> 206 gaatnttngg gaaaagggct tacaggacta gaaaccaaata angaaaanta atnntaangg 660
W--> 207 cnttatcntn aaaggtnata accnctccta tnatcccaac caatngnatt ccccaacnnc 720
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219 <223> OTHER INFORMATION: n = A,T,C or G
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224 ctgaagcgca cgtcccagaa ggtggacttg gcactgaaac agctgggaca catccgcgag 180
225 tacgaacagc gctgaaagt gctggagcgg gaggtccagc agtgtagccg cgtcctgggg 240
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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/483,672ADATE: 07/30/2000  
TIME: 17:04:25Input Set : A:\Pto.amc  
Output Set : N:\CRF3\07282000\I483672A.raw

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W--> 228 ggattttgct cctanantaa ggctcatctg ggctcggcc ccccaacctg gttggccttg 420
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W--> 235 ctttccctct nggganncg 799
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240 <213> ORGANISM: Homo sapien
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250 caaggacaag gccaccaggt gcgggggccc aagcccacat gatccttact ctatgagcaa 180
W--> 251 aatccctgtt gggggcttct ccttgaagtc cgccancagg gctcagcttt tggaccang 240
W--> 252 caggctcatg ggtgtgngc caactggggg ccncaacgca aaanggcna gggcctcngn 300
W--> 253 caccatccc angacgcggc tacactnctg gacctccnc tccaccatt tcatgcgctg 360
W--> 254 ttentacccg cgnatntgtc ccactgttt cngtgcncac tccancttct nggacgtgcg 420
W--> 255 ctacatacgc ccggantcnc nctcccgctt tgctccatc cagctnccan caacaaattt 480
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W--> 259 gggaanance ctcgncnctn ccccncttaa tccncccttg cnangnncnt ccccnntcc 720
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263 <210> SEQ ID NO: 10
264 <211> LENGTH: 789
265 <212> TYPE: DNA
266 <213> ORGANISM: Homo sapien
268 <220> FEATURE:
269 <221> NAME/KEY: misc_feature
270 <222> LOCATION: (1)...(789)
271 <223> OTHER INFORMATION: n = A,T,C or G
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276 agatcctgcc ctacacactg gcctccctct accaccggga gaagcagggt ttctgcccc 180
277 aataccgagg ggacactgga ggtgctagca gtgaggacag cctgatgacc agcttctgc 240
278 caggccctaa gcctggagct ccttcccta atggacacgt ggggtgctga ggcagtggcc 300
279 tgcctccacc tccacccgcg ctctgcgggg cctctgcctg tgatgtctcc gtacgtgtgg 360
W--> 280 tgggtgggtga gccaccgan gccaggggtg ttccgggccc gggcatctgc ctggacctgc 420
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**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

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VERIFICATION SUMMARY

DATE: 07/30/2000

PATENT APPLICATION: US/09/483,672A

TIME: 17:04:26

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07282000\I483672A.raw

L:23 M:270 C: Current Application Number differs, Wrong Format  
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:52 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:54 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:76 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:80 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:97 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:126 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:128 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:156 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:157 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:182 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:183 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:204 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:205 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:207 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:209 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:227 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:228 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:234 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8

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Input Set : A:\Pto.amc  
Output Set: N:\CRF3\07282000\I483672A.raw

L:235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8  
L:249 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9  
L:251 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9  
L:698 M:283 W: Missing Blank Line separator, <210> field identifier  
L:1467 M:283 W: Missing Blank Line separator, <400> field identifier  
L:10018 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:502  
L:10018 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:502  
L:10018 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:502  
L:10018 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:502  
L:10018 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:502  
L:10019 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:502  
L:10019 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:502  
L:10019 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:502  
L:10019 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:502  
M:340 Repeated in SeqNo=502  
L:10020 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:502  
L:10020 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:502  
L:10020 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:502  
L:10020 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:502  
L:10021 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:502  
L:10021 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:502  
L:10021 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:502  
L:10021 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:502  
L:10023 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:502  
L:10023 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:502  
L:10023 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:502  
L:10023 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:502  
L:10031 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:503  
L:10031 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:503  
L:10031 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:503  
L:10031 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:503  
L:10031 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:503  
L:10032 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:503  
L:10032 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:503  
L:10032 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:503  
L:10032 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:503  
M:340 Repeated in SeqNo=503  
L:10033 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:503  
L:10033 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:503  
L:10033 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:503  
L:10033 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:503  
L:10035 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:503  
L:10035 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:503  
L:10035 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:503  
L:10035 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:503  
L:10036 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:503  
L:10036 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:503  
L:10036 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:503  
L:10036 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:503



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Input Set : A:\Pto.amc

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L:10037 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:503  
L:10037 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:503  
L:10037 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:503  
L:10037 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:503  
L:10107 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:508  
L:10107 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:508  
L:10107 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:508  
L:10107 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:508  
L:10107 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:508